Figure 1 Nucleotide sequence and translation of HCQAS17

	· 10 30 50	
1	50	60
C1	70 90 110	
61 1	TAAGGTCTCTGAGATCCTTGCACTAGCTACATCCTCAGGGTAGGAGGAAGATGGCTTCCA M A S R	120 4
	130 150 170	
121	GAAGCATGCGGCTGCTCCTATTGCTGAGCTGCCTGGCCAAAACAGGAGTCCTGGGTGATA	180
5	SMRLLLLSCLAKTGVLGDI	24
181	190 210 230 TCATCATGAGACCCAGCTGTGCTCCTGGATGGTTTTACCACAAGTCCAATTGCTATGGTT	240
25	I M R P S C A P G W F Y H K S N C Y G Y	44 44
	250 270 290	
241 45	ACTTCAGGAAGCTGAGGAACTGGTCTGATGCCGAGCTCGAGTGTCAGTCTTACGGAAACG	300
43		64
301	310 330 350 GAGCCCACCTGGCATCTATCCTGAGTTTAAAGGAAGCCAGCACCATAGCAGAGTACATAA	360
65	A H L A S I L S L K E A S T I A E Y I S	84
	370 390 410	
361	GTGGCTATCAGAGAAGCCAGCCGATATGGATTGGCCTGCACGACCCACAGAAGAGGCAGC	420
85	G Y Q R S Q P I W I G L H D P Q K R Q Q	104
121	430 450 470 AGTGGCAGTGGATTGATGGGGCCATGTATCTGTACAGATCCTGGTCTGGCAAGTCCATGG	400
105	W Q W I D G A M Y L Y R S W S G K S M G	480 124
	490 510 530	
181	GTGGGAACAAGCACTGTGCTGAGATGAGCTCCAATAACAACTTTTTAACTTGGAGCAGCA	540
L25	G N K H C A E M S S N N N F L T W S S N	144
541	550 570 590 ACGAATGCAACAAGCGCCAACACTTCCTGTGCAAGTACCGACCATAGAGCAAGAATCAAG	600
45	E C N K R Q H F L C K Y R P *	600 164
	610 630 650	
501	ATTCTGCTAACTCCTGCACAGCCCCGTCCTCTTCCTTTCTGCTAGCCTGGCTAAATCTGC	660
	670 690 710	
661	TCATTATTTCAGAGGGGAAACCTAGCAAACTAAGAGTGATAAGGGCCCTACTACACTGGC	720
	730 750 770	
721	TTTTTTAGGCTTAGAGACAĞAAACTTTAGCATTGGCCCAGTAGTGGCTTCTAGCTCTAAA	780
	790 810 830	
781	TGTTTGCCCCGCCATCCCTTTCCACAGTATCCTTCTTCCCTCCTCCCCTGTCTCTGGCTG	840
	FIGURE 1 1/2 10+2	

841	850 TCTCGAGCAGTCTAGAAGAGTGCATCT	870 CCAGCCTA	TGAAACAGCTG	890 GTCTTTGGCCATA	900
901	910 AGAAGTAAAGATTTGAAGACAGAAGGA	930 AGAAACTC	AGGAGTAAGCT	950 FCTAGACCCCTTCA	960
961	970 GCTTCTACACCCTTCTGCCCTCTCTCC	990 ATTGCCTG	: CACCCACCCC	LØ10 AGCCACTCAACTCC	1020
1021	1030 TGCTTGTTTTTCCTTTGGCCATAGGAA	050 GGTTTACC	1 AGTAGAATCCTT	.070 GCTAGGTTGATGT	1080
1081	1090 1 GGGCCATACATTCCTTTAATAAACCAT	110 TGTGTAC	1114		

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